

09/807258

Figure 1

SEQ ID NO: 7	MSSL-MISTAMKGYAPY-RQVRDGYIAQPHN	C	AYH	C	LKISSG	C	DTL	C	KENGATSGH	C	GHK
SEQ ID NO: 2	-----LALLEMTGYES-VRDGYIAQPHN	C	VYH	C	IPD---	C	DTL	C	KDNGGTGGH	C	GFK
SEQ ID NO: 4	-----SLALLEMTGYES-VRDGYIAQPHN	C	AHH	C	FPGSSG	C	DTL	C	KENGGTGGH	C	GFK
SEQ ID NO: 6	MNHLVMISLALLEMTGYESGVRDGYIAQPHN	C	VYH	C	FPGSPG	C	DTL	C	KENGASSGH	C	GFK

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SEQ ID NO: 7	SGHGSA	C	W	C	KDLPDKVGIIVHGEK	C	HR
SEQ ID NO: 2	LGHGIA	C	W	C	NALPDNVGIIVDGVK	C	HK
SEQ ID NO: 4	VGHGTA	C	W	C	NALPDNVGIIVDGVK	C	HR
SEQ ID NO: 6	EGHGLA	C	W	C	NDLPDKVGIIVGEK	C	HK

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FIGURE 2

SEQ ID NO:10	-----GRDAYIAQPEN	<input type="checkbox"/> VYE	<input type="checkbox"/> AQNSY	<input type="checkbox"/> NDL	<input type="checkbox"/> TKNGATSGY	<input type="checkbox"/> QWL GK
SEQ ID NO:09	MNYLVXISLALLMTGVESGRDAYIAQNYN	<input type="checkbox"/> VYH	<input type="checkbox"/> ALNPY	<input type="checkbox"/> NDL	<input type="checkbox"/> TKNGAKSGY	<input type="checkbox"/> QWFGS
	1					60

SEQ ID NO:10	YGNA	<input type="checkbox"/> W	<input type="checkbox"/> KDLPDNVPIRIPGK	<input type="checkbox"/> HF
SEQ ID NO:09	SGNA	<input type="checkbox"/> W	<input type="checkbox"/> IDLPDNPVPIKVP GK	<input type="checkbox"/> HR
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FIGURE 3

SEQ ID NO:17	-----DGYI-KRRDG	C	KVA	C	LIGNEG	C	DKE	C	KAYGGSYGY	C	WTW
SEQ ID NO:12	MKLLLLLIVSASMLIESLVNADGYI-RRKDG	C	KVA	C	LFGNDG	C	NKE	C	KAYGAYYGY	C	WTW
SEQ ID NO:14	MKLLLLLTISASMLIEGLVNADGYI-RGGDG	C	KVS	C	VINHVF	C	DNE	C	KAAGGSYGY	C	WAW
SEQ ID NO:16	MKIIIFLIVSSMLI-GVKTDNGYLLNKATG	C	KVW	C	VINNAS	C	NSE	C	KLRRGNNGYGY	C	YEW

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SEQ ID NO:17	GLA	C	W	C	EGLPDDKTWKSETNT	C	E---
SEQ ID NO:12	GLA	C	W	C	EGLPDDKTWKSETNT	C	GGKK
SEQ ID NO:14	GLA	C	W	C	EGLPADREWKYETNT	C	GGKK
SEQ ID NO:16	KLA	C	Y	C	EGAPKSELWAYATNK	C	NGKL

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